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NTDB id 1131 NGFG RS05740 WP 003695064.1    ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 257129 CVE23 RS02870 WP 024109635.1 MASRGVNKVILIGNLGQDPEIRYMPNGGAVANLTMATSDTWRDKQTGEQKERTEWHRVTLYGKLAEIAGEYLRKGAQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                      **!!!!**!*!!*!!**!*!!*!*!!*****!!!**!*!* *!***!*!!!!*********!!*!!*!!*!!* !!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1    EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS..QEAYQRPAQQSR...QPA.......   140
NTDB id 1103 NMB RS07590 WP 002212976.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQS..QEAYQRPAQQSR...QPA.......   140
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQ.......GSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 257129 CVE23 RS02870 WP 024109635.1 EGQLRTRKWQDQSGQDRYTTEVVV.DINGSMQMLGNRSGA.............QNPANSWGQPQQPEAPSHS.........   138
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGMNVPAQQGSWGQPQQPAKQ.HQ.........   149
consensus                                   !!****!!*!** !**!*******      **!!**!  * *  *   *      * ***** ****   *
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NTDB id 1131 NGFG RS05740 WP 003695064.1    PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1103 NMB RS07590 WP 002212976.1     SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF.................   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF.................   180
NTDB id 257129 CVE23 RS02870 WP 024109635.1 ....GMPQNSAPQ.AQRSAGNEPP.MDFDDDIPFAALGLSIPNHALYAIS   182
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ....PMQQSAPQQ.YSQPQYNEPP.MDFDDDIPF.................   177
consensus                                      ** **     ***  ***!****!!!!!!
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X non conserved
X similar
X ≥ 50% conserved